CRF Errors Corrected by the STIC Systems Branch

~~ENTERED~~

Serial Number: __09/900,497__

CRF Processing Date: __1/22/2002__
Edited by: _____
Verified by: _____ (STIC staff)

- ☐ Changed a file from non-ASCII to ASCII

- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.

- ☐ Edited a format error in the Current Application Data section, specifically:
_____

- ☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other _____.

- ☐ Added the mandatory heading and subheadings for "Current Application Data".

- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.

- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically:
_____

- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:
_____

- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:
_____

- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.

- ☐ Inserted colons after headings/subheadings. Headings edited included:
_____

- ☐ Deleted extra, invalid, headings used by an applicant, specifically:
_____

- ☑ Deleted: ☑ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as_____ .

- ☐ Inserted mandatory headings, specifically: _____

- ☐ Corrected an obvious error in the response, specifically:
_____

- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.

- ☐ Corrected an error in the Number of Sequences field, specifically:
_____

- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.

- ☐ Deleted *ending* stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____

- ☐ Other:
_____
_____
_____

*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

3/1/95

OIPE

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/900,497

DATE: 01/22/2002
TIME: 19:36:22

Input Set : A:\PTO.AMC.txt
Output Set: N:\CRF3\01222002\I900497.raw

```
 4 <110> APPLICANT: Allen, Keith D.
 6 <120> TITLE OF INVENTION: TRANSGENIC MICE CONTAINING NPY6-R
 7        NEUROPEPTIDE RECEPTOR GENE DISRUPTIONS
10 <130> FILE REFERENCE: R-639
12 <140> CURRENT APPLICATION NUMBER: US 09/900,497
13 <141> CURRENT FILING DATE: 2001-07-06
15 <150> PRIOR APPLICATION NUMBER: US 60/216,260
16 <151> PRIOR FILING DATE: 2000-07-06
18 <150> PRIOR APPLICATION NUMBER: US 60/221,474
19 <151> PRIOR FILING DATE: 2000-07-27
21 <160> NUMBER OF SEQ ID NOS: 3
23 <170> SOFTWARE: FastSEQ for Windows Version 4.0
25 <210> SEQ ID NO: 1
26 <211> LENGTH: 2281
27 <212> TYPE: DNA
28 <213> ORGANISM: Mus musculus
30 <400> SEQUENCE: 1
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32 aaaggggatg gaaatatata cttgtactgc cttagatagt caccaggatg ttgttacagt 120
33 cttcgtttac tgcttctgaa gcctatactg atagaattaa taaaatactg agagagagag 180
34 agagggacag agagagagag ggggagagag agagagagag agagagagag agagagagag 240
35 agagagagag agaagagaag aaaacaaggt saagccatct gcttaactta tgtccacatt 300
36 ctctcaagag cattgtccta tttgtagaat tatctatatt gttaagaatc atctccattg 360
37 ttaagatttt gtgggctgga gatccagctc tgttgataaa gtgcttgcct aacatgcatg 420
38 aagtcctagg ttctattccc aaggctacat aaaaccttgt gttgtgatga atgcctgtaa 480
39 tcccagtacg cagcaaggag agacaaggag gatcagaagc ttaaggacat cattttgtac 540
40 atagtgagtt tgaggaaagc tgaggttaca tggaactctc tctctctcaa aaacaaaaca 600
41 aaacaaaaca aaaccttcta ctaatattct ggattctgtt tgatttttag gatctcaaga 660
42 gcatgctgac gtcatttatg tgtttccatc agatacagac agagatcata aacatttaac 720
43 tcattgatta tatgttgaga gttgtccctc aagaaccaat ggccaaacat ccactgagga 780
44 tacacggaag cttagaaaat ctctaattaa aatcctgaca taatggaagt gctcacaaac 840
45 cagccaacac ctaataaaac cagtggcaag agcaacaact cggcattttt ctactttgaa 900
46 tcctgccaac ccccttttct agccatactc ttgctactca tagcatatac tgtgatccta 960
47 atcatgggca tttttggaaa cctctctctt atcatcatca tctttaagaa acagagagaa 1020
48 gctcaaaatg ttaccaacat actgattgcc aacctgtccc tctctgacat cttggtgtgt 1080
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55 ctcctctttt ctacatcatt atttatgctc cagtattttg tccctctggg tttcattctt 1500
56 atctgctacc tgaagatcgt tctctgcctc cgaaaaagaa ctaggcaggt ggacaggaga 1560
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57 aaggaaaata agagccgtct caatgagaac aagagggtaa atgtgatgtt gatttccatc 1620
58 gtagtgactt ttggagcctg ctggttgccc ttgaacattt tcaatgtcat cttcgactgg 1680
59 tatcatgaga tgctgatgag ctgccaccac gacctggtat ttgtagtttg ccacttgatt 1740
60 gctatggttt ctacttgcat aaatcctctc ttttatggat ttctcaacaa aaacttccag 1800
61 aaggatctaa tgatgcttat tcaccactgt tggtgtggtg aacctcagga aagttatgaa 1860
62 aatattgcca tgtctactat gcacacagat gaatccaagg gatcattaaa actggctcac 1920
63 ataccaacag gcatatagaa actggtaagc aaaatcaaag cccttctgtt atgaaagaaa 1980
64 gagaagaaat agtatggaat agggcaaggt gcagaggaag ccagacttaa acacataata 2040
65 tctttgggcc cagttttgct ttaagttaag catgtctact ccattcagcc atagaacaca 2100
66 cagagattta tccctaccct ttcttttttt cctttggaag aataataact taaacaacct 2160
67 agacatcatt actgaggaag agaacaaaaa tgagagagca tacaaggaca gcagagatgt 2220
68 ctggggtaca aaattcacgt tattcgctgg aatagctaga aagttattag ttgtgctgca 2280
69 g                                                                 2281
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72 <211> LENGTH: 200
73 <212> TYPE: DNA
74 <213> ORGANISM: Artificial Sequence
76 <220> FEATURE:
77 <223> OTHER INFORMATION: Targeting Vector
79 <400> SEQUENCE: 2
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81 tcataaacat ttaactcatt gattatatgt tgagagttgt ccctcaagaa ccaatggcca 120
82 aacatccact gaggatacac ggaagcttag aaaatctcta attaaaatcc tgacataatg 180
83 gaagtgctca caaaccagcc                                              200
85 <210> SEQ ID NO: 3
86 <211> LENGTH: 200
87 <212> TYPE: DNA
88 <213> ORGANISM: Artificial Sequence
90 <220> FEATURE:
91 <223> OTHER INFORMATION: Targeting Vector
93 <400> SEQUENCE: 3
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95 tacactctga tggaccactg ggtatttggg aacactatgt gtaaactcac ttcctacgtg 120
96 caaagtgtct cagtttctgt gtccatattc tcccttgtgt tgattgctat tgaacgatat 180
97 cagctgattg tgaacccccg                                              200
```

VERIFICATION SUMMARY
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